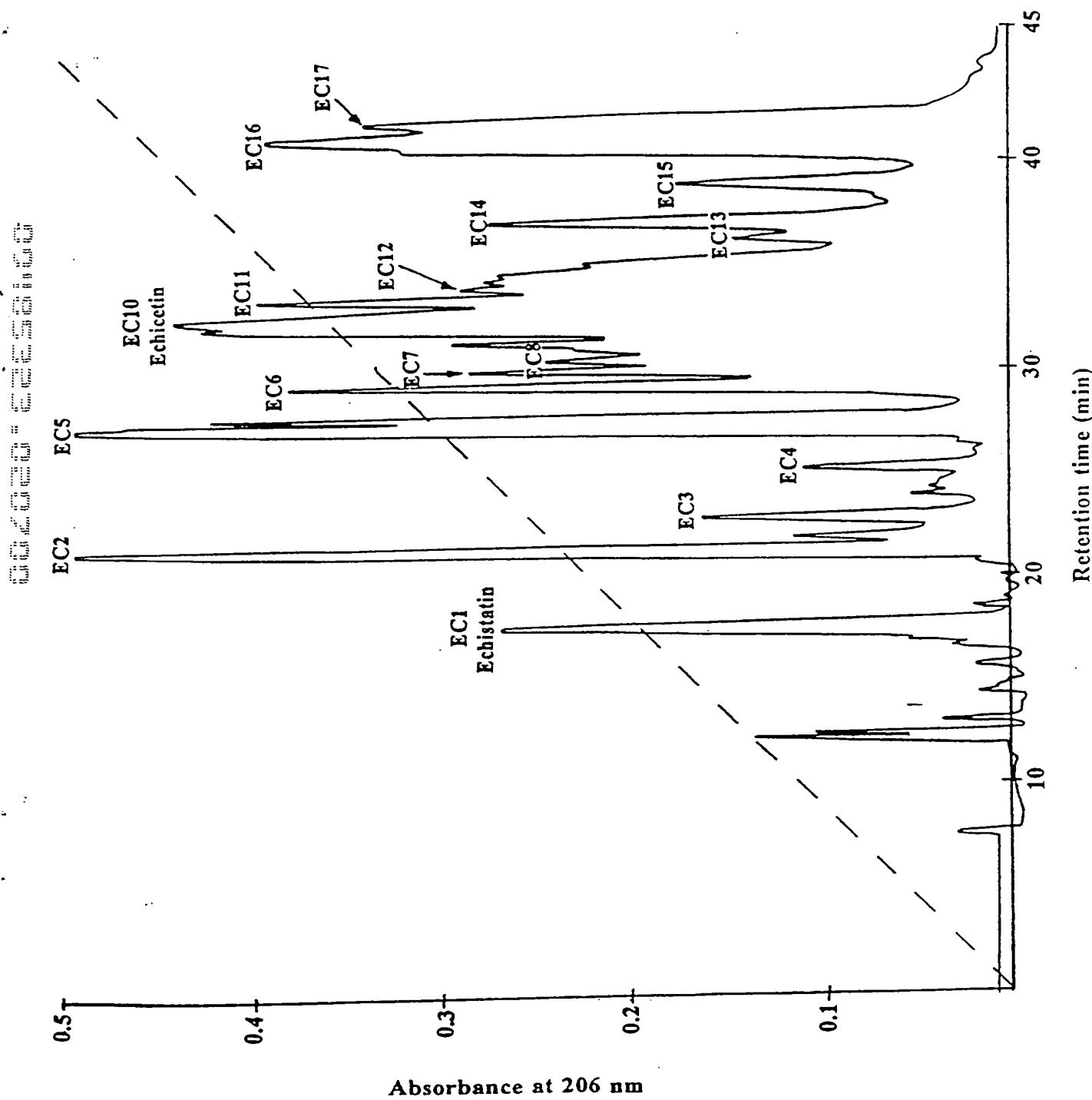
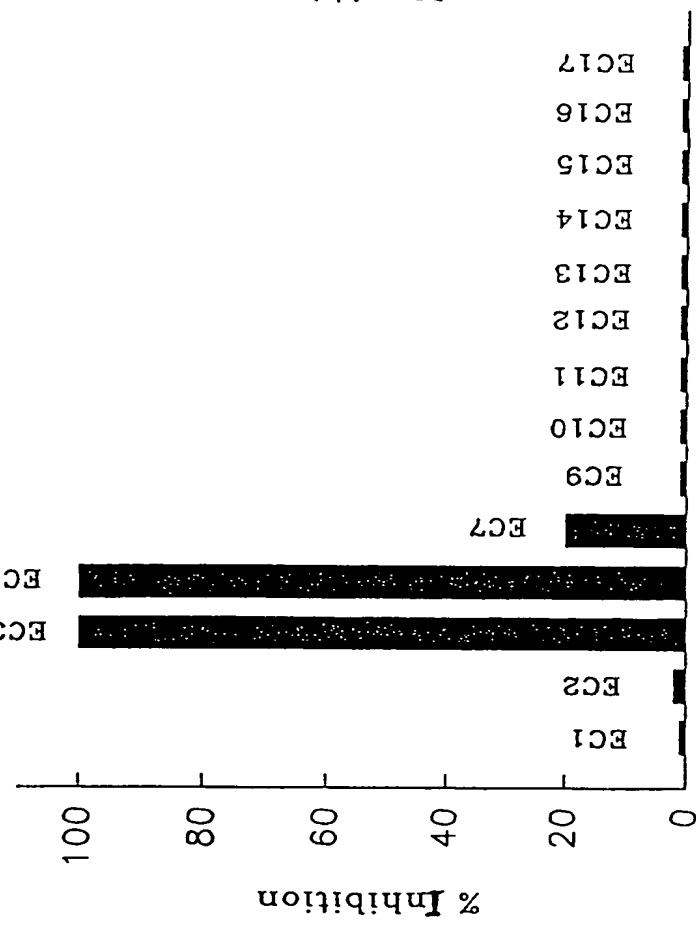
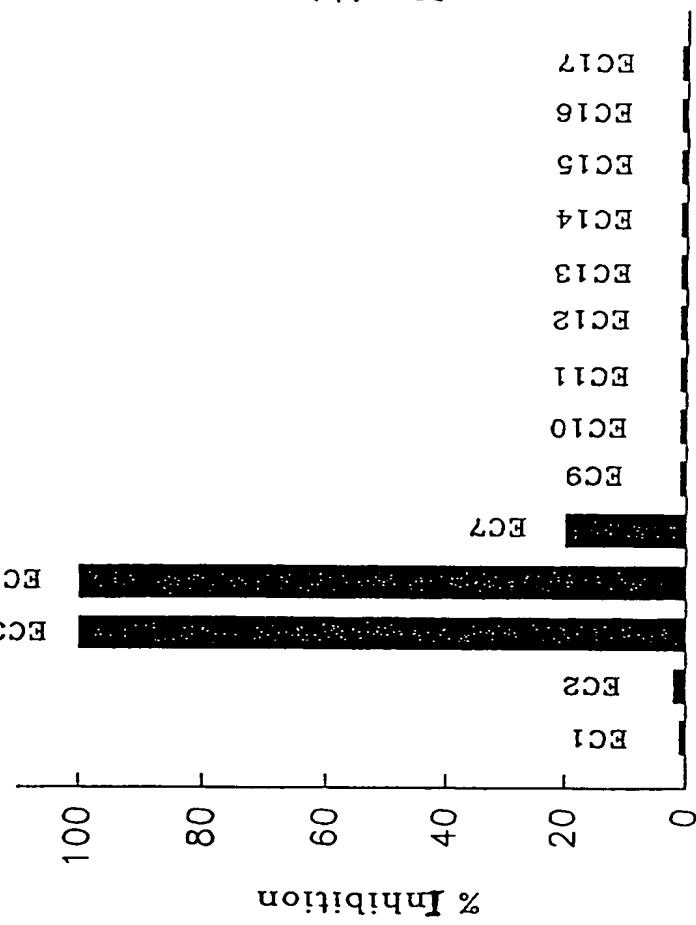


Fig. 1



2/12

Fig.  
2AFig.  
2B

3/12

Absorbance at 206 nm

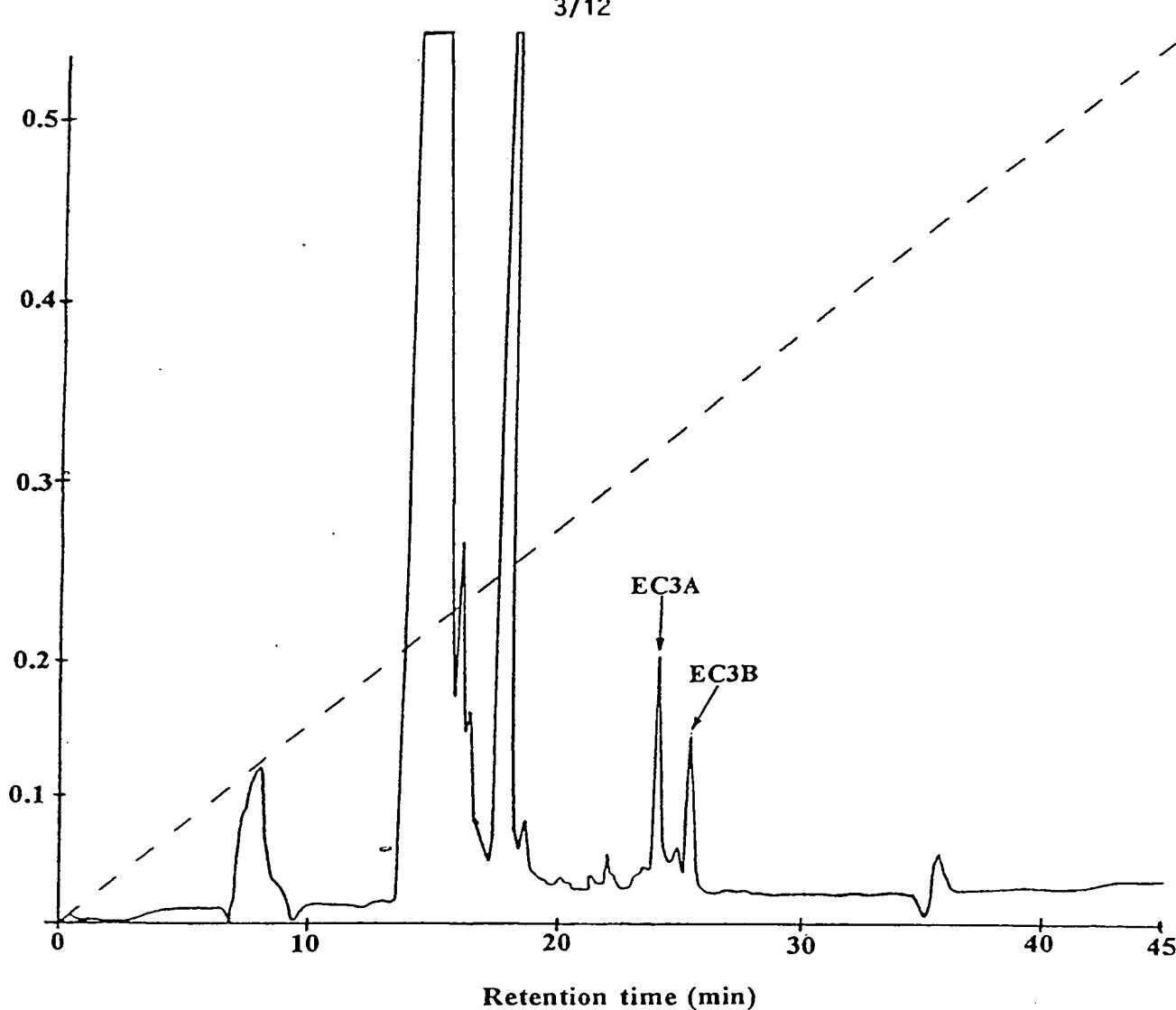


Fig. 3

1 NSVHPCCDPVKCEPREGEHCISGPCCRNCYFLRAGTVCKRAVGDDVDDYCSGITPDCFRNRYKGKED  
 |----- N-terminal sequencing -----|  
 |-----|-----|-----|-----|-----|  
 |----- K3 -----||-- R K1b V -|  
 |----- K1a -----|-- N K2 I -|----- K4 -----|

1 NSVHPCCDPVKCEPREGEHCISGPCCRNCKFLNAGTICKRAMLDGGLNDCTGISTDCFRNRYKGKED  
 |----- N-terminal sequencing -----|  
 |-----|-----|-----|-----|-----|  
 |----- T K7 -----||-- K6 -----|K----- K10a -- T -----|  
 |T ----- K4 -----|----- K10b -- S -----|

T

Fig. 4A

Fig. 4B

<u>Eristostatin</u>	QEEPCATGPCARROKFKRAGKVCRVA <u>RGD</u> WNDDYOTGKSODOPRNPWNG
<u>Echistatin</u>	ECESGPCORNCFLKEGTICKRA <u>RGD</u> DMDDYCNQGKTPDOPRNPWNGPAT
<u>Flavoridin</u>	GEECPDGSPPSNPCCDDAATCKLRLPGAOQCDGICCDQCRFKKTKGICRIA <u>RGD</u> FPDDRCTGLSNDOPRNWNDL
<u>Kistrin</u>	GKECPDSSPENPCCDDATCKLRLPGAOQCGEGGLCCEQCKFSRAGKICRIP <u>RGD</u> MPDDDRCTGQSADOPRYH
<u>EC3A</u>	NSVHPCCDPVKCEPREGEHCTISGPCCORNCFLRAGTVQKRA <u>VGD</u> DVDDYCSGSGITPDOPRNRYKGKED
<u>EC3B</u>	NSVHPCCDPVKCEPREGEHCTISGPCCORNCFLNAGTICKRA <u>MLD</u> GLNDYCTGKSSDOPRNRYKGKED
<u>Le3 (disintegrin domain)</u>	...NSGNPCCDPVTICPRRGEHCVSGKCCORNCFLRAGTVQKRA <u>VGD</u> DMDDYCTGCISSDOPRNPYXKD

Fig. 5

6/12

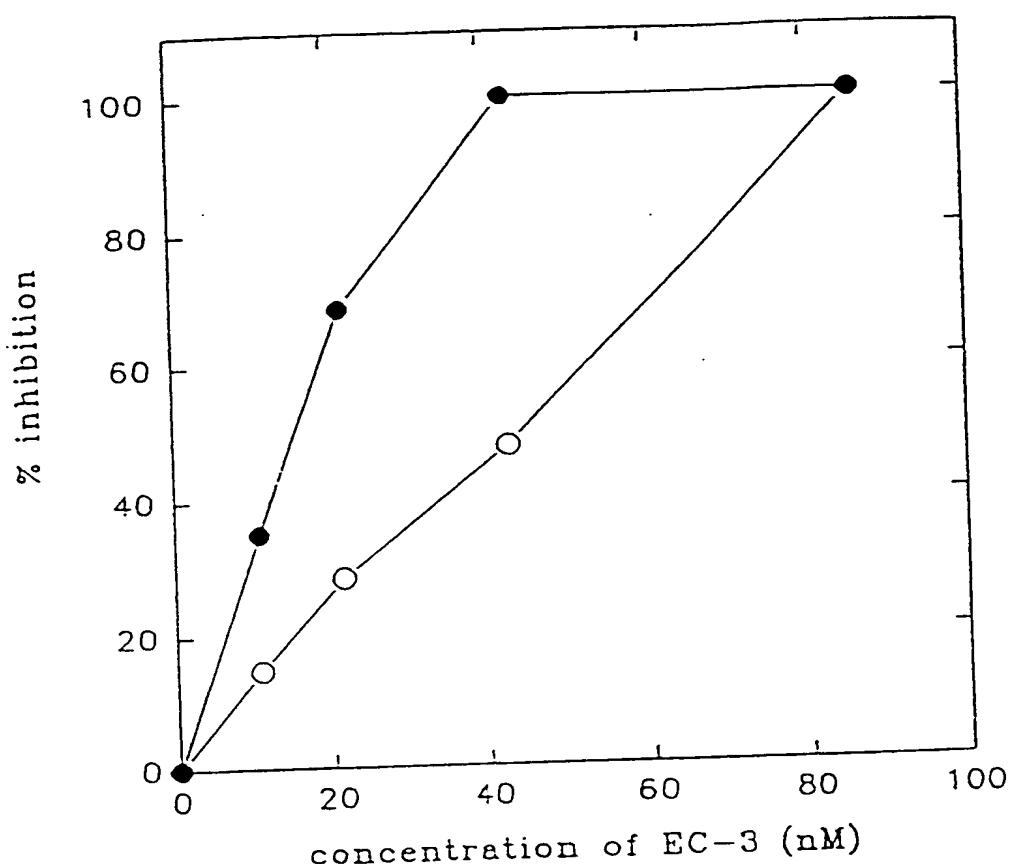


Fig. 6

7/12

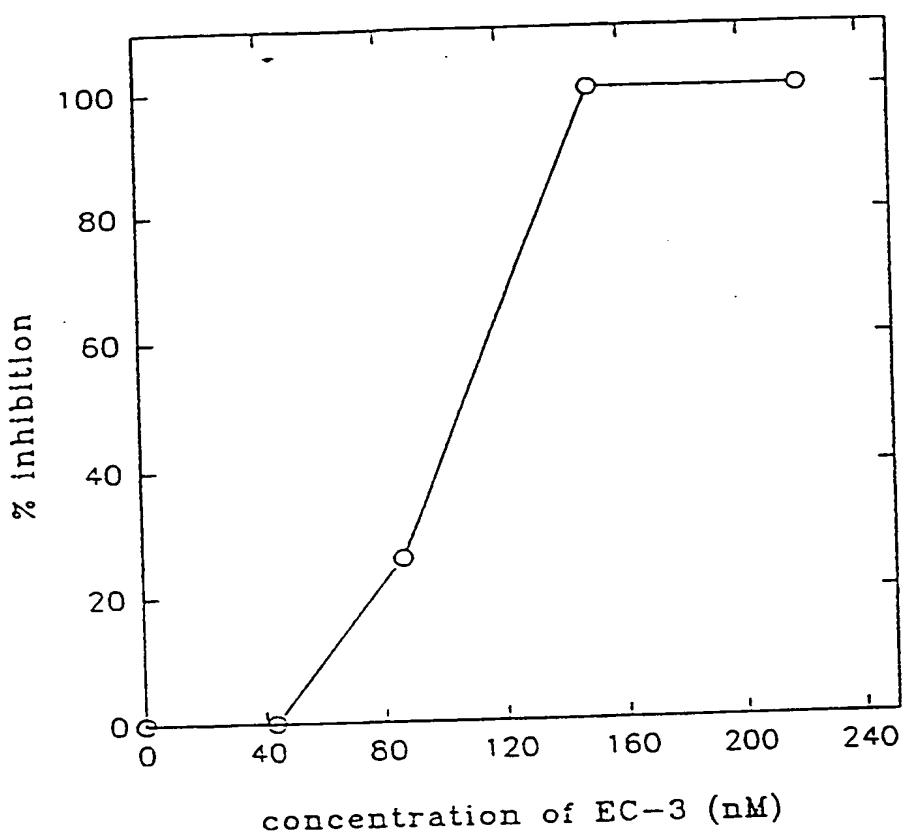
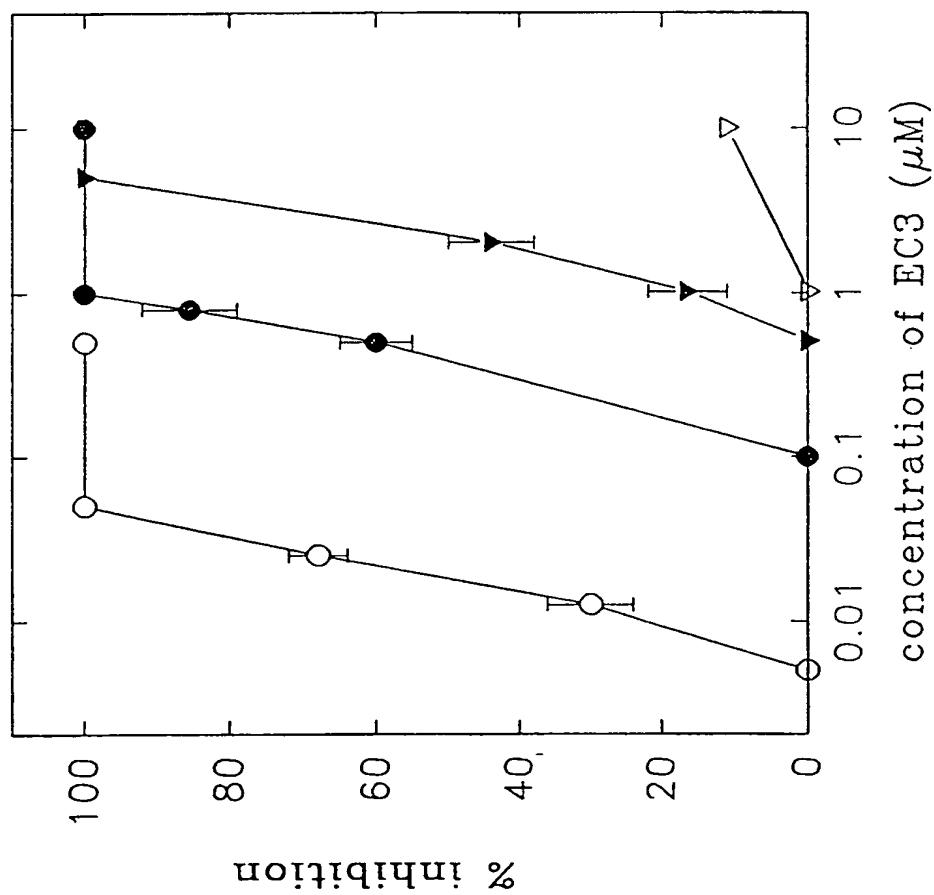


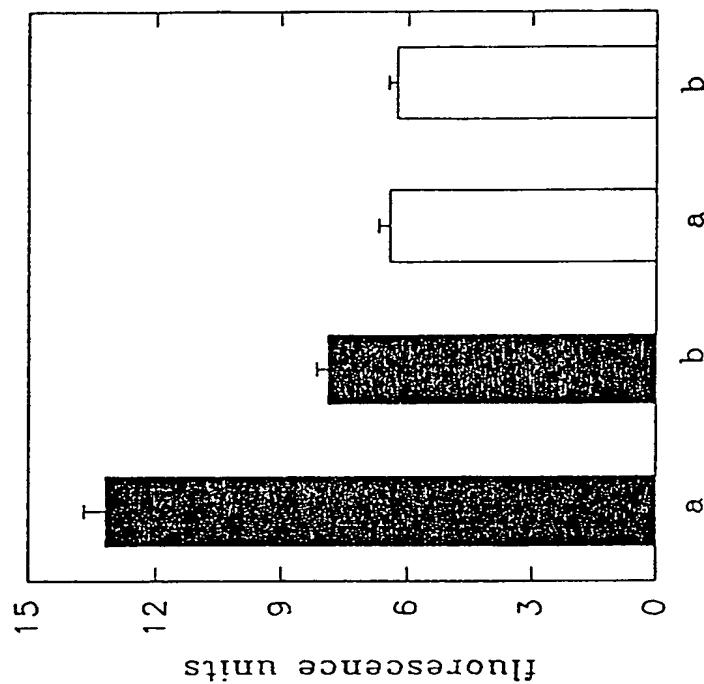
Fig. 7

Fig. 8



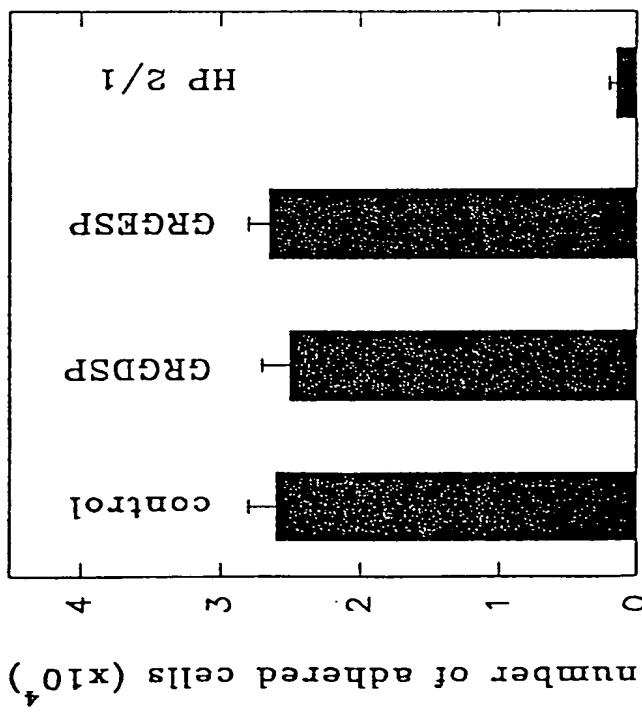
9/12

Fig. 9B



HP 2/1

Fig. 9A



10/12

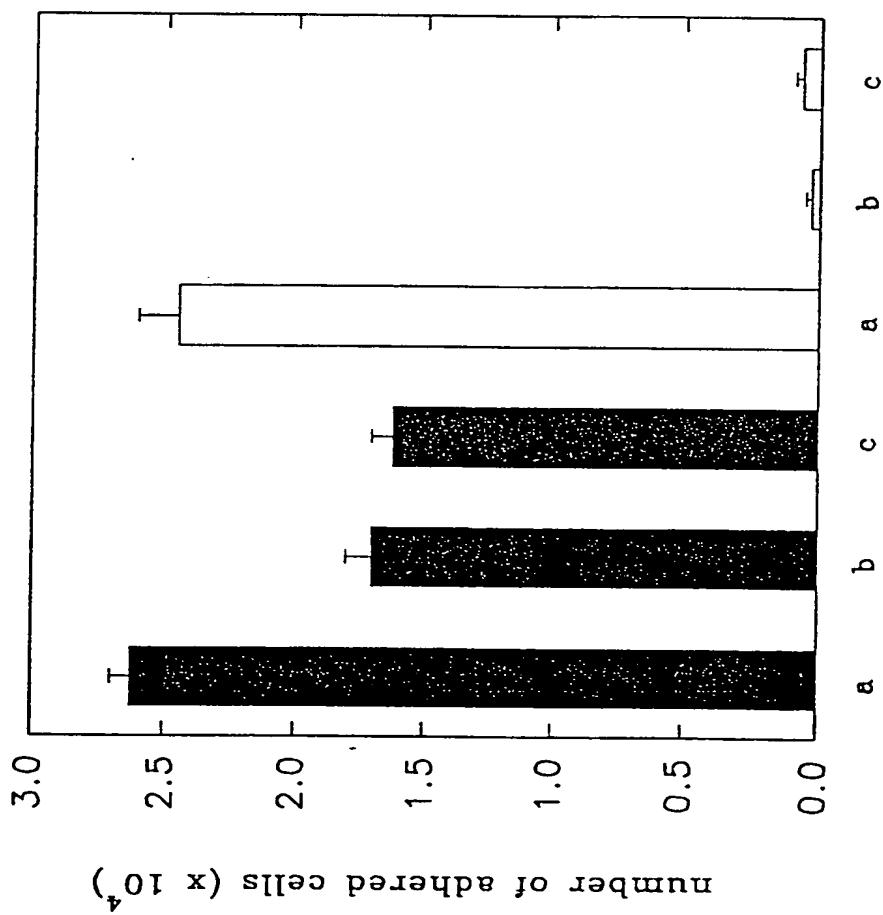
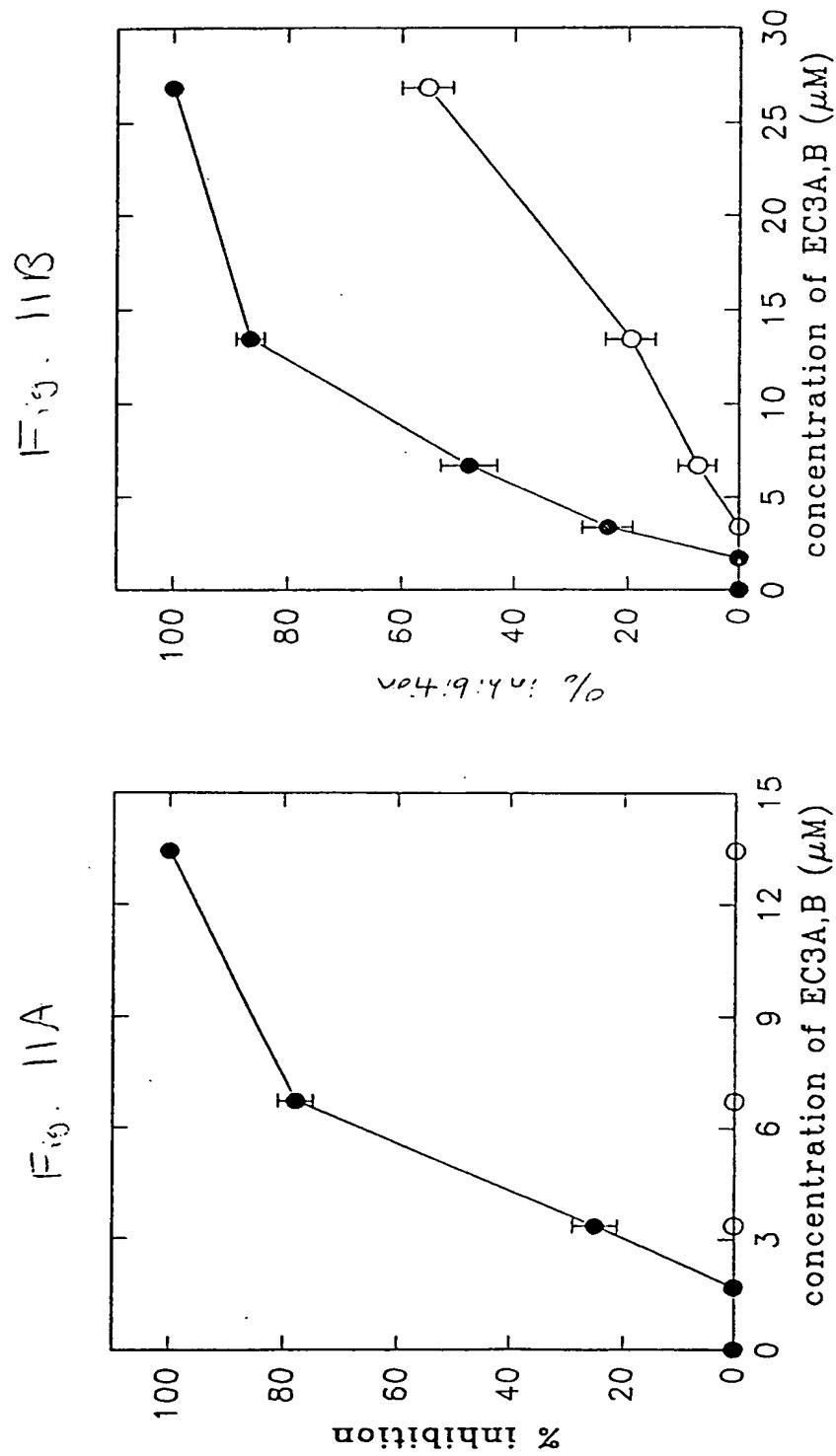


Fig. 10

11/12



12/12

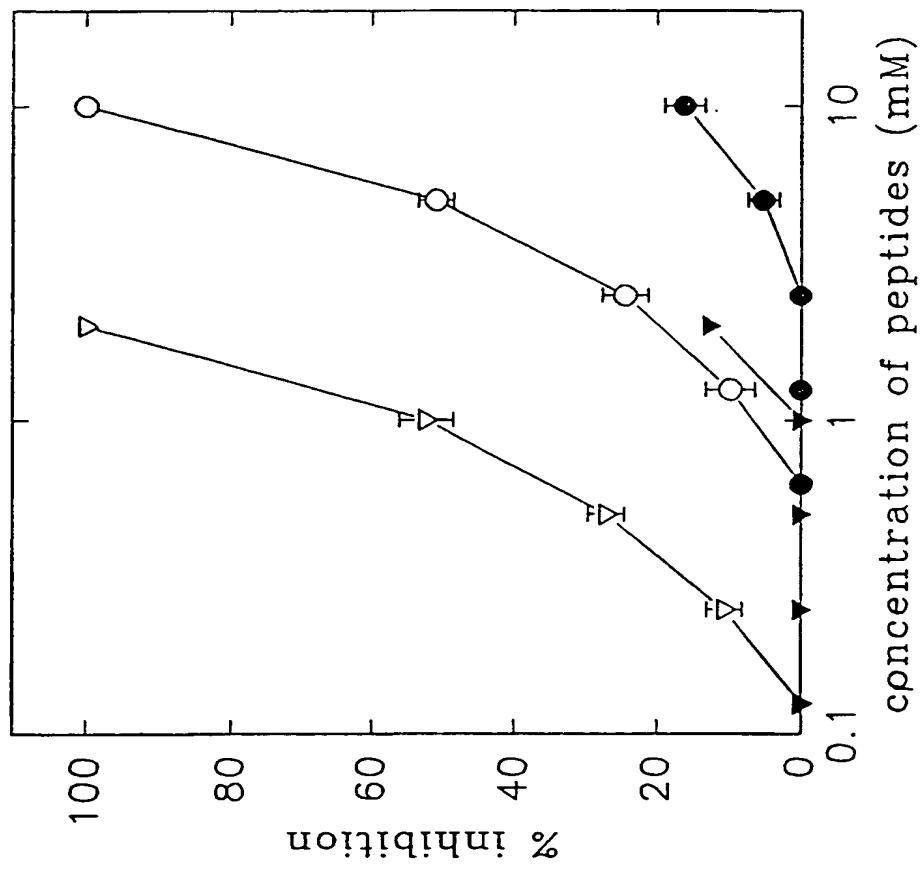


Fig. 12